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SEQUENCE LISTING

<110> KAWAOKA, Akiyoshi
EBINUMA, Hiroyasu

<120> TRANSCRIPTION FACTOR CONTROLLING PHENYLPROPANOID
BIOSYNTHESIS PATHWAY

<130> 4859-0027-0

<140> 09/282,146

<141> 1999-03-31

<150> JP 10-125171

<151> 1998-03-31

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<170> PatentIn Ver. 2.1

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| Thr Ala His Glu Gly Arg Leu Tyr Cys Lys His His His Ile Gln Leu | | | | | | | | | | | | | | | | | | | | |
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| Ile Lys Glu Lys Gly Asn Leu Ser Lys Leu Glu Gly Asp His Glu Met | | | | | | | | | | | | | | | | | | | | |
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SEQUENCE LENGTH: 200
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 MOLECULE TYPE: protein

SEQUENCE

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SEQUENCE LISTING

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SEQUENCE LENGTH: 988
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA to mRNA
ORIGINAL SOURCE

ORGANISM: *Nicotiana tobacum*
Tissue tepey: leaf

FEATURE

NAME/KEY: CDS
LOCATION: 100..702
IDENTIFICATION METHOD: E

NAME/KEY: domain

LOCATION: 127..282

IDENTIFICATION METHOD: S

ANOTHER INFORMATION: LIM domain which is a kind of the zing finger motif

NAME/KEY: domain

LOCATION: 427..582

IDENTIFICATION METHOD: S

ANOTHER INFORMATION: LIM domain which is a kind of the zing finger motif

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